

Fig. 14

GAATTCGGCACGAGGGATCTGGATGGCATCTACTTCGTATGACTATTGCAGAGTGCCCAT 60  
     M A S T S Y D Y C R V P M 13  
 GGAAGACGGGGATAAGCGCTGTAAGCTTCTGCTGGGATAGGAATTCTGGTGCTCCTGAT 120  
 E D G D K R C K L L L G I G I L V L L I 33  
 CATCGTGATTCTGGGGTGCCCTTGATTATCTTCACCATCAAGGCCAACAGCGAGGCCTG 180  
 I V I L G V P L I I F T I K A N S E A C 53  
 CCGGACGGCCCTTCGGGCAGTGATGGAGTGTCGCAATGTCACCCATCTCCTGCAACAAGA 240  
 R D G L R A V M E C R N V T H L L Q Q E 73  
 GCTGACCGAGGCCAGAGGGCTTTCAGGATGTGGAGGCCCGCCGACCTGCAACCA 300  
 L T E A Q K G F Q D V E A Q A A T C N H 93  
 CACTGTGATGGCCCTAATGGCTTCCCTGGATGCAGAGAAGGCCCAAGGACAAAAGAAAGT 360  
     T V M A L M A S L D A E K A Q G Q K K V 113  
 GGAGGAGCTTGAGGGAGAGATCACTACATTAAACCATAAGCTTCAGGACGCGTCTGCAGA 420  
 E E L E G E I T T L N H K L Q D A S A E 133  
 GGTGAGCGACTGAGAAGAGAAAACCAGGTCTTAAGCGTGAGAATCGCGGACAAAGATA 480  
 V E R L R R E N Q V L S V R I A D K K Y 153  
 CTACCCAGCTCCAGGACTCCAGCTCCGCTGCGGGGCCCCAGCTGCTGATTGTGCTGCT 540  
 Y P S S Q D S S A A A P Q L L I V L L 173  
 GGGCCTCAGCGCTCTGCTGCAGTGAGATCCCAGGAAGCTGGCACATCTTGGAAGGTCCGT 600  
     G L S A L L Q \* (SEQ ID NO:26) 180  
 CCTGCTCGGCTTTTCGCTTGAACATTCCCTTGATCTCATCAGTCTGTAGCGGGTCATGGG 660  
 GCAACACGGTTAGCGGGGAGAGCAGCGGGTAGCCGGAGAGGGCCCTCTGGAGCAGGTCTG 720  
 GAGGGCCCATGGGGCAGTCCCTGGGTGTGGGGACACAGTCGGGTGACCCAGGCTGTCTC 780  
 CCTCGAGAGCCCTCCCTCCGACAAATGAGTCCCCCTCTTGTCTCCACCCCTGAGATTGGG 840  
 CATGGGGTGGGGTGGGGGCATGTGCTGCCCTGTTGTTATGGGTTTTCGCGGGGG 900  
 GGTGCTTTTTCGCGGTCTTTGAGCTCCAAAAAATAAACACTTCCTTTGAGGGAGAG 960  
 CACACCTTAAAAAATAAAAAAATAAAAAAATAAATTCGGGGCGCGCCA 1014  
 (SEQ ID NO:16)



Fig.15

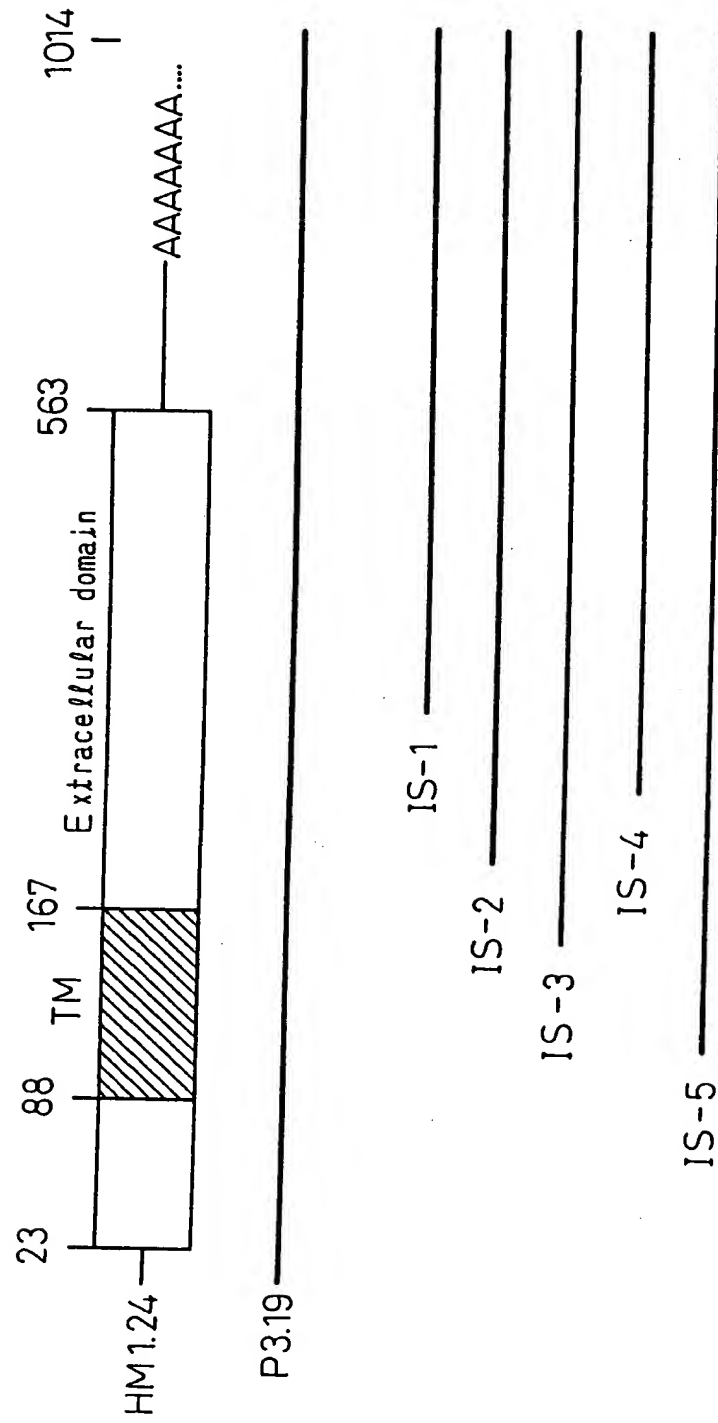


Fig.16

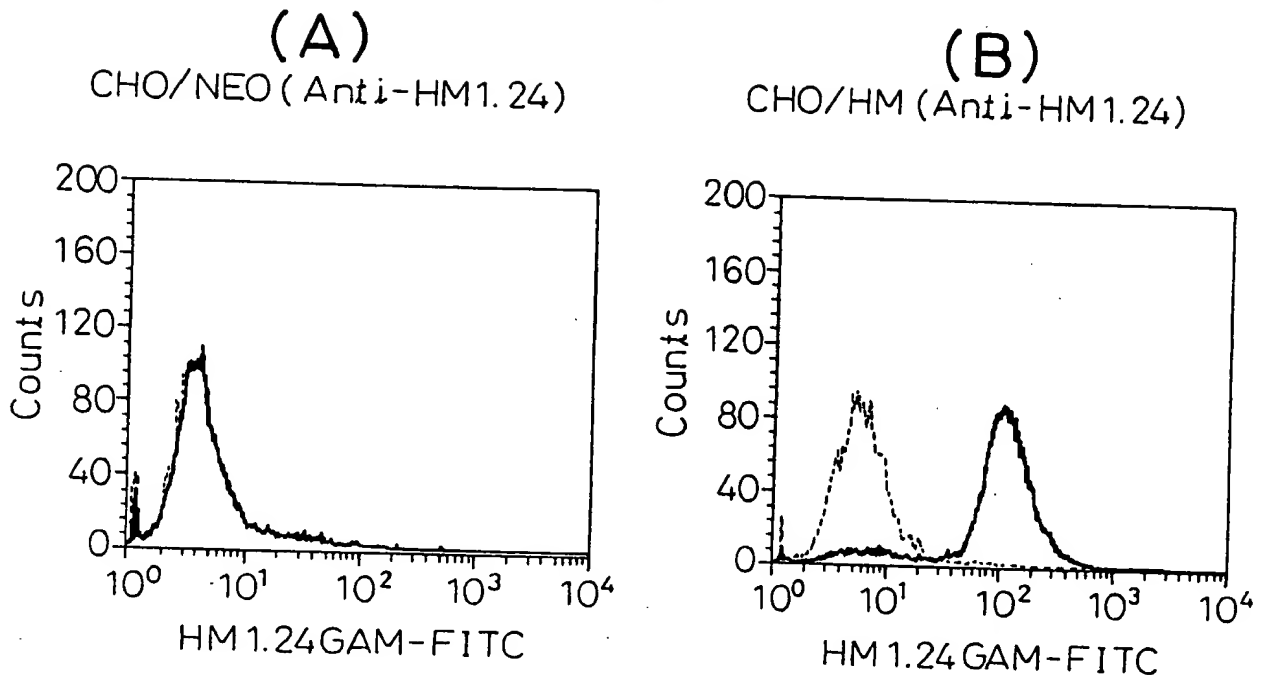
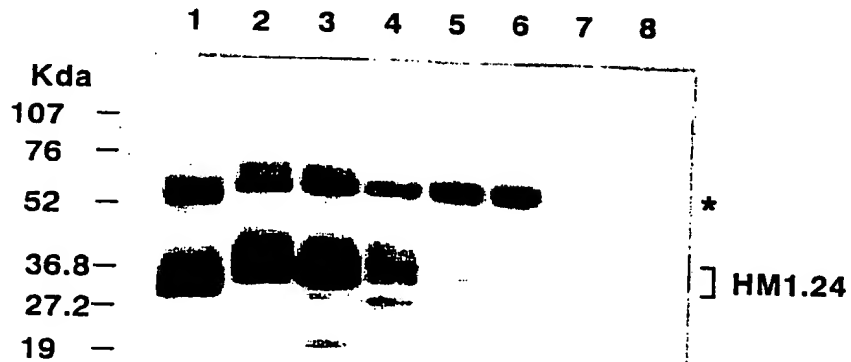


Fig.17



LANE 1: KPMM2 (EQUIVALENT TO  $5 \times 10^5$  CELLS)

LANE 2: RPMI8226 ( $25 \times 10^5$  CELLS)

LANE 3: U266 ( $25 \times 10^5$  CELLS)

LANE 4: CHO/HM ( $5 \times 10^5$  CELLS)

LANE 5: CHO/NEO ( $5 \times 10^5$  CELLS)

LANE 6: NONE

LANE 7: KPMM2 ( $5 \times 10^5$  CELLS)

Fig.18

